

## SEQUENCE LISTING

## MAR 2 4 2005 OFFICE OF PETITIONS

35

PANGALOS, Menelas
NEEFS, Jean-Marc
PEETERS, Danielle

<120>	CLONING	AND	CHARACTERISATION	OF	NOVEL	MAMMALIAN	PEPTIDASES

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<141> 2001-01-12

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<151> 1999-07-14

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Ser Cys His Arg Thr Glu Glu Asn Val Thr Gly Glu Gln Gly Gly Pro 130 135 140

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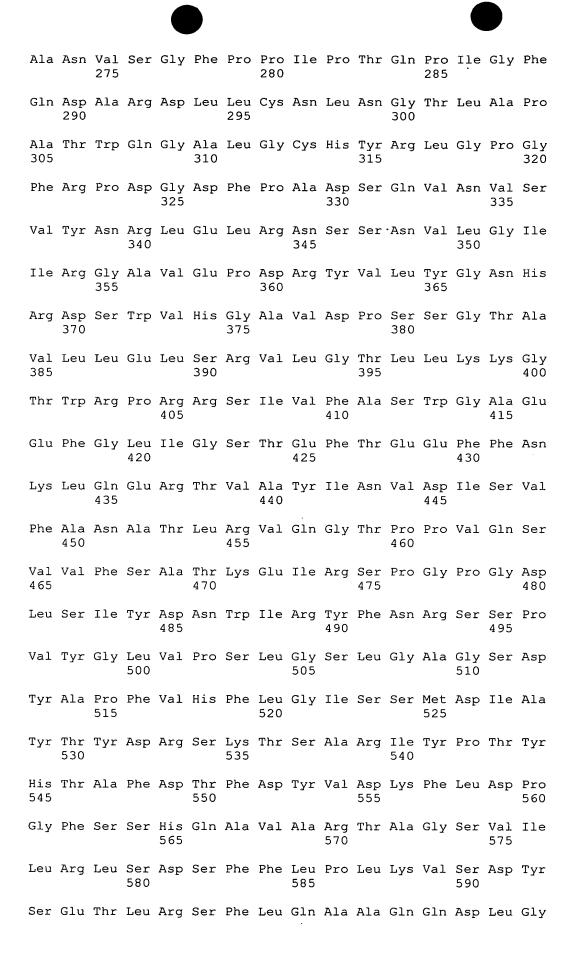
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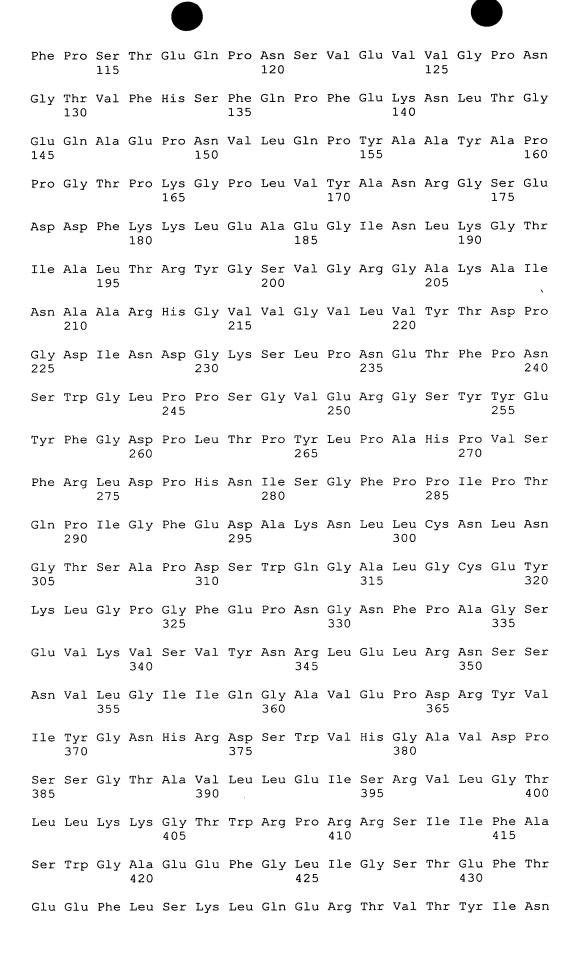
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Ser Gly Leu Asp Thr Ala Lys Thr Tyr Glu Tyr Thr Val Leu Leu Ser 100 105 110



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Pro Leu Val Thr Ala Val Glu Lys Phe Lys Ala Ala Ala Ala Ala Ala Glu Arg Met Val Asn Asn Gln His Bro Arg Ala Phe G75  Ala Pro Asn Thr Ala Ser Val Ala Thr Phe Pro Gly Leu Ala Asn G90  Tyr Ala Arg Ala Gln Glu Ile Asn Ser Gly Ala Glu Ala Trp Ala	Asn	Val		Asp	Tyr	Ser	Glu		Leu	Gln	Ser	Phe		Gln	Ala	Ala
Asn Gln His Ile Leu Thr Leu Gln Lys Ser Ser Pro Asp Pro Leu 655  Val Arg Met Val Asn Asp Gln Leu Met Leu Leu Glu Arg Ala Phe 660  Asn Pro Arg Ala Phe Pro Glu Glu Arg Tyr Tyr Ser His Val Leu 675  Ala Pro Asn Thr Ala Ser Val Ala Thr Phe Pro Gly Leu Ala Asn 690  Tyr Ala Arg Ala Gln Glu Ile Asn Ser Gly Ala Glu Ala Trp Ala	Gln		Asn	Leu	Gly	Ala		Leu	Glu	Ser	His		Ile	Ser	Leu	Gly
Val Arg Met Val Asn Asp Gln Leu Met Leu Leu Glu Arg Ala Phe 660  Asn Pro Arg Ala Phe Pro Glu Glu Arg Tyr Tyr Ser His Val Leu 675  Ala Pro Asn Thr Ala Ser Val Ala Thr Phe Pro Gly Leu Ala Asn 690  Tyr Ala Arg Ala Gln Glu Ile Asn Ser Gly Ala Glu Ala Trp Ala		Leu	Val	Thr	Ala		Glu	Lys	Pḥe	Lys		Ala	Ala	Ala	Ala	Leu 640
Asn Pro Arg Ala Phe Pro Glu Glu Arg Tyr Tyr Ser His Val Leu 675  Ala Pro Asn Thr Ala Ser Val Ala Thr Phe Pro Gly Leu Ala Asn 690  Tyr Ala Arg Ala Gln Glu Ile Asn Ser Gly Ala Glu Ala Trp Ala	Asn	Gln	His	Ile		Thr	Leu	Gln	Lys		Ser	Pro	Asp	Pro	Leu 655	Gln
Ala Pro Asn Thr Ala Ser Val Ala Thr Phe Pro Gly Leu Ala Asr 690 695 700  Tyr Ala Arg Ala Gln Glu Ile Asn Ser Gly Ala Glu Ala Trp Ala	Val	Arg	Met		Asn	Asp	Gln	Leu		Leu	Leu	Glu	Arg		Phe	Leu
690 695 700  Tyr Ala Arg Ala Gln Glu Ile Asn Ser Gly Ala Glu Ala Trp Ala	Asn	Pro		Ala	Phe	Pro	Glu		Arg	Tyr	Tyr	Ser		Val	Leu	Trp
	Ala		Asn	Thr	Ala	Ser		Ala	Thr	Phe	Pro		Leu	Ala	Asn	Ala
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<212> PRT

<213> Homo sapiens

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Tyr Gly Gly Val Gly Arg Gly Ala Lys

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<212> PRT

<213> Homo sapiens

<400> 38

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Pro Ala Asp 35

<210> 39

<211> 20

<212> PRT

<213> Homo sapiens

<400> 39

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Ser Pro Ala Gln

20

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Pro Leu	Thr	Met 20	Trp	Thr	Ser	Phe	Trp 25	Thr	Arg	Ala	Ser	Ala 30	Ala	Ile		
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gtgagcg ggggctg tctggat gcccacc	gtet a	agcct gctgt	ggto	ga go ca to	ccago	catat	tgo	etged	ctgc	acco	ccago	gee d	cctgo	ctct	gc	120 180
gtgagcg ggggctg tctggat gcccacc	gtet agteg agge getet d	agcct gctgt	ggto	ga go ca to	ccago	catat	tgo	etged	ctgc	acco	ccago	gee d	cctgo	ctct	gc	120 180
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gtgageg ggggetg tetggat geecace <210> <211> <212> <213> <400> Gly Glu	gtct agtgg agge control of the second	sapi	cectoctects  Lens  Ser 5	ga go	cceto ccago ggtto	cetet ecet <u>e</u> etet <u>e</u>	tgc	Pro 10	etge ecac etce	acco	ccago ccago ggcca	gcc o	Cys	Ser	gc	120 180

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<400> 46

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<210> 47

<211> 3110

<212> DNA

<213> Homo sapiens

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3060

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3110

<210> 48

<211> 740

<212> PRT

<213> Homo sapiens

<400> 48

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Pro Leu Lys Glu Thr Thr Ser Val Arg Tyr His Gln Ser Ile Arg 35 40 45

Trp Lys Leu Val Ser Glu Met Lys Ala Glu Asn Ile Lys Ser Phe Leu 50 55 60

Arg Ser Phe Thr Lys Leu Pro His Leu Ala Gly Thr Glu Gln Asn Phe 65 70 . 75 80

Leu Leu Ala Lys Lys Ile Gln Thr Gln Trp Lys Lys Phe Gly Leu Asp 85 90 95

Ser Ala Lys Leu Val His Tyr Asp Val Leu Leu Ser Tyr Pro Asn Glu 100 105 110

Thr Asn Ala Asn Tyr Ile Ser Ile Val Asp Glu His Glu Thr Glu Ile 115 120 125

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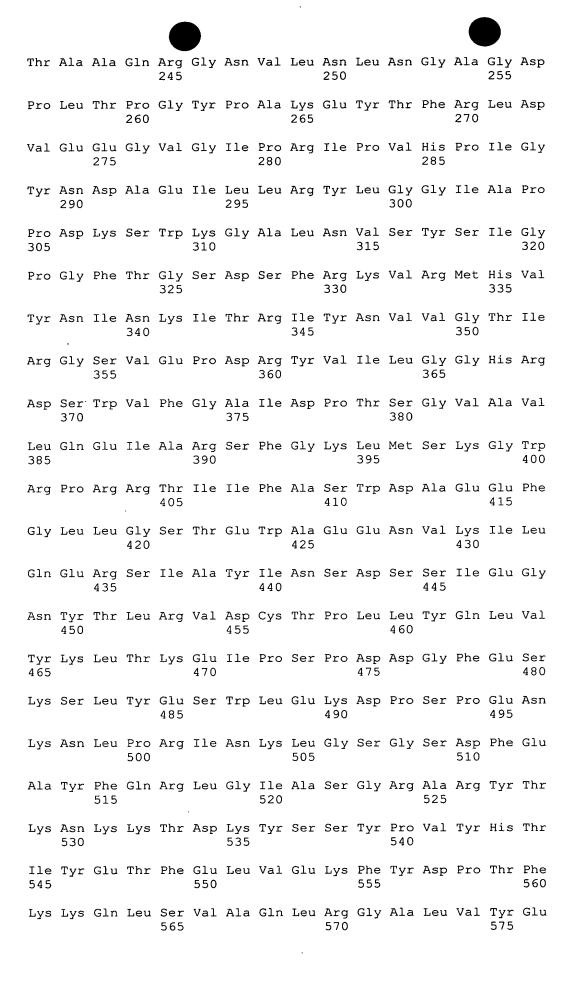
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Ala Arg Tyr Gly Lys Ile Phe Arg Gly Asn Lys Val Lys Asn Ala Met 195 200 205

Leu Ala Gly Ala Ile Gly Ile Ile Leu Tyr Ser Asp Pro Ala Asp Tyr 210 215 220

Phe Ala Pro Glu Val Gln Pro Tyr Pro Lys Gly Trp Asn Leu Pro Gly 225 230 235 240



Leu Val Asp Ser Lys Ile Ile Pro Phe Asn Ile Gln Asp Tyr Ala Glu 580 585 590

Ala Leu Lys Asn Tyr Ala Ala Ser Ile Tyr Asn Leu Ser Lys Lys His 595 600 605

Asp Gln Gln Leu Thr Asp His Gly Val Ser Phe Asp Ser Leu Phe Ser 610 620

Ala Val Lys Asn Phe Ser Glu Ala Ala Ser Asp Phe His Lys Arg Leu 625 630 635 640

Ile Gln Val Asp Leu Asn Asn Pro Ile Ala Val Arg Met Met Asn Asp 645 650 655

Gln Leu Met Leu Leu Glu Arg Ala Phe Ile Asp Pro Leu Gly Leu Pro 660 665 670

Gly Lys Leu Phe Tyr Arg His Ile Ile Phe Ala Pro Ser Ser His Asn 675 680 685 .

Lys Tyr Ala Gly Glu Ser Phe Pro Gly Ile Tyr Asp Ala Ile Phe Asp 690 695 . 700

Ile Glu Asn Lys Ala Asn Ser Arg Leu Ala Trp Lys Glu Val Lys 705 710 715 720

His Ile Ser Ile Ala Ala Phe Thr Ile Gln Ala Ala Ala Gly Thr Leu 725 730 735

Lys Glu Val Leu 740

<210> 49

<211> 1884

<212> DNA

<213> Homo sapiens

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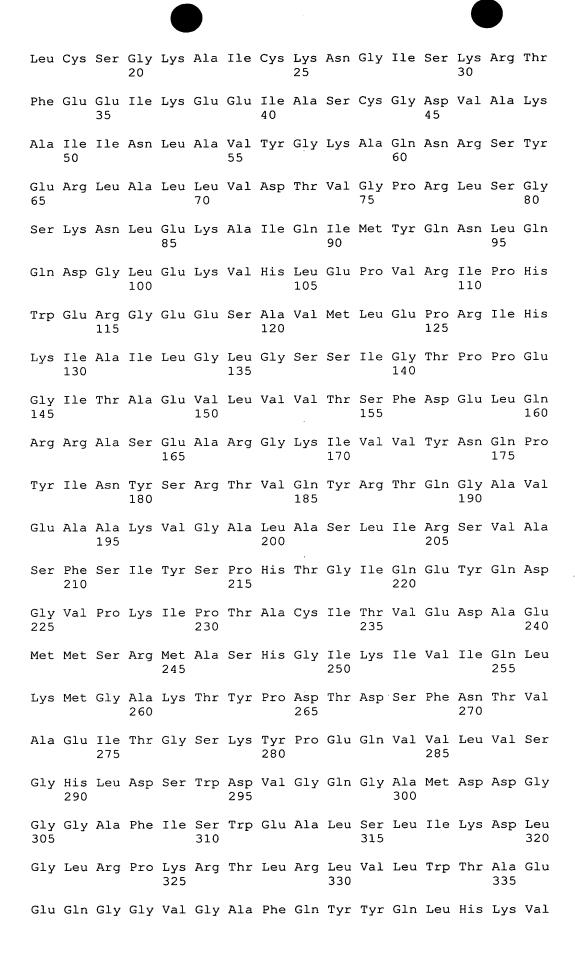
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<212> PRT

<213> Homo sapiens

<400> 50

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Leu Pro Thr Gly Leu Gln Phe Thr Gly Ser Glu Lys Ala Arg Ala Ile 370 375 380

Met Glu Glu Val Met Ser Leu Leu Gln Pro Leu Asn Ile Thr Gln Val 385 390 395 400

Leu Ser His Gly Glu Gly Thr Asp Ile Asn Phe Trp Ile Gln Ala Gly
405 410 415

Val Pro Gly Ala Ser Leu Leu Asp Asp Leu Tyr Lys Tyr Phe Phe 420 425 430

His His Ser His Gly Asp Thr Met Thr Val Met Asp Pro Lys Gln Met 435 440 445

Asn Val Ala Ala Val Trp Ala Val Val Ser Tyr Val Val Ala Asp 450 455 460

Met Glu Glu Met Leu Pro Arg Ser 465 470

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<210> 51

<211> 750

<212> PRT

<213> Homo sapiens

<400> 51

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Phe Leu Cly Phe Leu Phe Gly Trp Phe Ile Lys Ser Ser Asn Glu 35 40 45

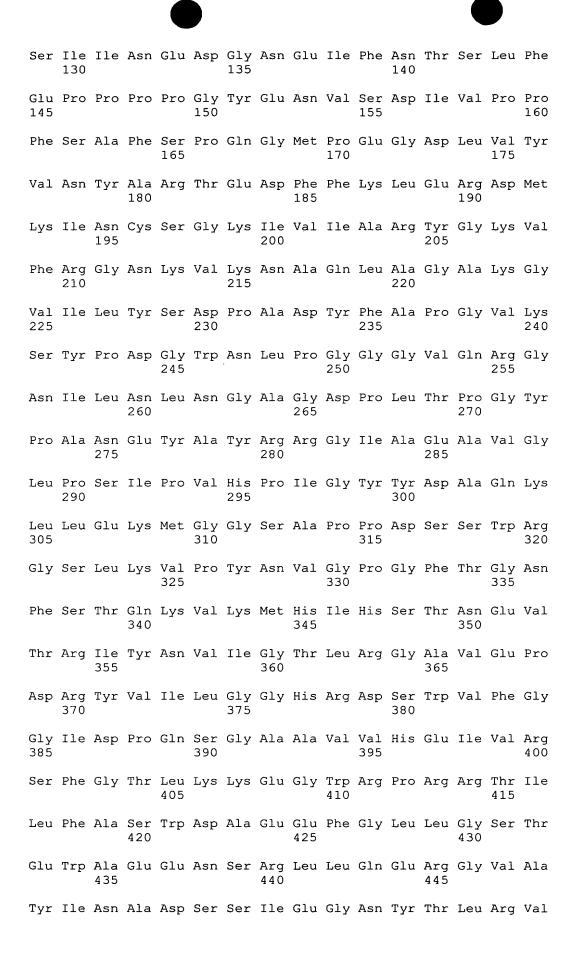
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Leu Lys Ala Glu Asn Ile Lys Lys Phe Leu His Asn Phe Thr Gln Ile 65 70 75 80

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Gln Ser Gln Trp Lys Glu Phe Gly Leu Asp Ser Val Glu Leu Ala His 100 105 110

Tyr Asp Val Leu Ser Tyr Pro Asn Lys Thr His Pro Asn Tyr Ile 115 120 125



450 455 460 Asp Cys Thr Pro Leu Met Tyr Ser Leu Val His Asn Leu Thr Lys Glu 470 475 Leu Lys Ser Pro Asp Glu Gly Phe Glu Gly Lys Ser Leu Tyr Glu Ser Trp Thr Lys Lys Ser Pro Ser Pro Glu Phe Ser Gly Met Pro Arg Ile 505 Ser Lys Leu Gly Ser Gly Asn Asp Phe Glu Val Phe Phe Gln Arg Leu Gly Ile Ala Ser Gly Arg Ala Arg Tyr Thr Lys Asn Trp Glu Thr Asn Lys Phe Ser Gly Tyr Pro Leu Tyr His Ser Val Tyr Glu Thr Tyr Glu 545 550 Leu Val Glu Lys Phe Tyr Asp Pro Met Phe Lys Tyr His Leu Thr Val Ala Gln Val Arg Gly Gly Met Val Phe Glu Leu Ala Asn Ser Ile Val 580 Leu Pro Phe Asp Cys Arg Asp Tyr Ala Val Val Leu Arg Lys Tyr Ala 600 Asp Lys Ile Tyr Ser Ile Ser Met Lys His Pro Gln Glu Met Lys Thr 610 615 Tyr Ser Val Ser Phe Asp Ser Leu Phe Ser Ala Val Lys Asn Phe Thr 630 635 Glu Ile Ala Ser Lys Phe Ser Glu Arg Leu Gln Asp Phe Asp Lys Ser 645 650 655 Asn Pro Ile Val Leu Arg Met Met Asn Asp Gln Leu Met Phe Leu Glu 665 Arg Ala Phe Ile Asp Pro Leu Gly Leu Pro Asp Arg Pro Phe Tyr Arg 680 His Val Ile Tyr Ala Pro Ser Ser His Asn Lys Tyr Ala Gly Glu Ser

Phe Thr Val Gln Ala Ala Ala Glu Thr Leu Ser Glu Val Ala 740 745 750

Phe Pro Gly Ile Tyr Asp Ala Leu Phe Asp Ile Glu Ser Lys Val Asp

Pro Ser Lys Ala Trp Gly Glu Val Lys Arg Gln Ile Tyr Val Ala Ala

<210> 52

<211> 265

PRT

<213> Saccharomyces cerevisiae

<400> 52

<212>

Thr Lys His Thr Val Ala Thr Val Gly Val Pro Tyr Lys Val Gly Lys
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Asp Thr Lys His Gly Asp Pro Asp Asn Ile Val Ala Leu Gly Ala His 50 55 60

Ser Asp Ser Val Glu Glu Gly Pro Gly Ile Asn Asp Asp Gly Ser Gly 65 70 75 80

Thr Ile Ser Leu Leu Asn Val Ala Lys Gln Leu Thr His Phe Lys Ile 85 90 95

Asn Asn Lys Val Arg Phe Ala Trp Trp Ala Ala Glu Glu Glu Gly Leu 100 105 110

Leu Gly Ser Asn Phe Tyr Ala Tyr Asn Leu Thr Lys Glu Glu Asn Ser 115 120 125

Lys Ile Arg Val Phe Met Asp Tyr Asp Met Met Ala Ser Pro Asn Tyr 130 135 140

Glu Tyr Glu Ile Tyr Asp Ala Asn Asn Lys Glu Asn Pro Lys Gly Ser 145 150 155 160

Glu Glu Leu Lys Asn Leu Tyr Val Asp Tyr Tyr Lys Ala His His Leu 165 170 175

Asn Tyr Thr Leu Val Pro Phe Asp Gly Arg Ser Asp Tyr Val Gly Phe 180 185 190

Ile Asn Asn Gly Ile Pro Ala Gly Gly Ile Ala Thr Gly Ala Glu Lys 195 200 205

Asn Asn Val Asn Asn Gly Lys Val Leu Asp Arg Cys Tyr His Gln Leu 210 215 220

Cys Asp Asp Val Ser Asn Leu Ser Trp Asp Ala Phe Ile Thr Asn Thr

Lys Leu Ile Ala His Ser Val Ala Thr Tyr Ala Asp Ser Phe Glu Gly 245 250 255

Phe Pro Lys Arg Glu Thr Gln Lys His 260 265 <211> 268

<212> PRT

<213> Vibrio cholerae

<400> 53

Gln Ile Thr Asn Thr Ile Arg Ala Leu Ser Ser Phe Asn Asn Arg Phe  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Tyr Thr Thr Ala Ser Gly Ala Gln Ala Ser Asp Trp Leu Ala Asn Glu 20 25 30

Trp Arg Ser Leu Ile Ser Ser Leu Pro Gly Ser Arg Ile Glu Gln Ile 35 40 45

Lys His Ser Gly Tyr Asn Gln Lys Ser Val Val Leu Thr Ile Gln Gly 50 55 60

Ser Glu Lys Pro Asp Glu Trp Val Ile Val Gly Gly His Leu Asp Ser 65 70 75 80

Asp Asp Ala Ser Gly Ile Ala Ser Leu Ser Glu Ile Ile Arg Val Leu 100 105 110

Arg Asp Asn Asn Phe Arg Pro Lys Arg Ser Ala Ala Leu Met Ala Tyr 115 120 125

Ala Ala Glu Glu Val Gly Leu Arg Gly Ser Gln Asp Pro Ala Asn Gln 130 135 140

Tyr Lys Ala Gln Gly Lys Lys Val Val Ser Val Leu Gln Leu Asp Met 145 150 155 160

Thr Asn Tyr Arg Gly Ser Ala Glu Asp Ile Val Phe Ile Thr Asp Tyr 165 170 175

Thr Asp Ser Asn Leu Thr Gln Phe Leu Thr Thr Leu Ile Asp Glu Tyr 180 185 190

Leu Pro Glu Leu Thr Tyr Gly Tyr Asp Arg Cys Gly Tyr Ala Cys Ser 195 200 205

Asp His Ala Ser Trp His Lys Ala Gly Phe Ser Ala Ala Met Pro Phe 210 215 220

Glu Ser Lys Phe Lys Asp Tyr Asn Pro Lys Ile His Thr Ser Gln Asp 225 230 235 240

Thr Leu Ala Asn Ser Asp Pro Thr Gly Asn His Ala Val Thr Phe Thr 245 250 255

Lys Leu Gly Leu Ala Tyr Val Ile Glu Met Ala Asn 260 265

<210> 54

<211> 268

<212> PRT

<213> Aeromonas proteolytica

<400> 54

Gln Ile Thr Gly Thr Ile Ser Ser Leu Glu Ser Phe Thr Asn Arg Phe  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Tyr Thr Thr Ser Gly Ala Gln Ala Ser Asp Trp Ile Ala Ser Glu 20 25 30

Trp Gln Ala Leu Ser Ala Ser Leu Pro Asn Ala Ser Val Lys Gln Val 35 40 45

Ser His Ser Gly Tyr Asn Gln Lys Ser Val Val Met Thr Ile Thr Gly 50 55 60

Ser Glu Ala Pro Asp Glu Trp Ile Val Ile Gly Gly His Leu Asp Ser 65 70 75 80

Thr Ile Gly Ser His Thr Asn Glu Gln Ser Val Ala Pro Gly Ala Asp 85 90 95

Asp Asp Ala Ser Gly Ile Ala Ala Val Thr Glu Val Ile Arg Val Leu 100 105 110

Ser Glu Asn Asn Phe Gln Pro Lys Arg Ser Ile Ala Phe Met Ala Tyr 115 120 125

Ala Ala Glu Glu Val Gly Leu Arg Gly Ser Gln Asp Leu Ala Asn Gln
130 135 140

Tyr Lys Ser Glu Gly Lys Asn Val Val Ser Ala Leu Gln Leu Asp Met 145 150 155 160

Thr Asn Tyr Lys Gly Ser Ala Gln Asp Val Val Phe Ile Thr Asp Tyr 165 170 175

Thr Asp Ser Asn Phe Thr Gln Tyr Leu Thr Gln Leu Met Asp Glu Tyr 180 185 190

Leu Pro Ser Leu Thr Tyr Gly Phe Asp Thr Cys Gly Tyr Ala Cys Ser 195 200 205

Asp His Ala Ser Trp His Asn Ala Gly Tyr Pro Ala Ala Met Pro Phe 210 215 220

Glu Ser Lys Phe Asn Asp Tyr Asn Pro Arg Ile His Thr Thr Gln Asp 225 230 235 240

Thr Leu Ala Asn Ser Asp Pro Thr Gly Ser His Ala Lys Lys Phe Thr 245 250 255

Gln Leu Gly Leu Ala Tyr Ala Ile Glu Met Gly Ser 260 265 <210> 55

<211> 263

<212> PRT

<213> Streptomyces griseus

<400> 55

Asn Asn Gly Gly Asn Arg Ala His Gly Arg Pro Gly Tyr Lys Ala Ser  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Val Asp Tyr Val Lys Ala Lys Leu Asp Ala Ala Gly Tyr Thr Thr Thr 20 25 30

Leu Gln Gln Phe Thr Ser Gly Gly Ala Thr Gly Tyr Asn Leu Ile Ala 35 40 45

Asn Trp Pro Gly Gly Asp Pro Asn Lys Val Leu Met Ala Gly Ala His 50 55 60

Leu Asp Ser Val Ser Ser Gly Ala Gly Ile Asn Asp Asn Gly Ser Gly 65 70 75 80

Ser Ala Ala Val Leu Glu Thr Ala Leu Ala Val Ser Arg Ala Gly Tyr 85 90 95

Gln Pro Asp Lys His Leu Arg Phe Ala Trp Trp Gly Ala Glu Glu Leu 100 105 110

Gly Leu Ile Gly Ser Lys Phe Tyr Val Asn Asn Leu Pro Ser Ala Asp 115 120 125

Arg Ser Lys Leu Ala Gly Tyr Leu Asn Phe Asp Met Ile Gly Ser Pro 130 135 140

Asn Pro Gly Tyr Phe Val Tyr Asp Asp Asp Pro Val Ile Glu Lys Thr 145 150 155 160

Phe Lys Asn Tyr Phe Ala Gly Leu Asn Val Pro Thr Glu Ile Glu Thr 165 170 175

Glu Gly Asp Gly Arg Ser Asp His Ala Pro Phe Lys Asn Val Gly Val 180 185 190

Pro Val Gly Gly Leu Phe Thr Gly Ala Gly Tyr Thr Lys Ser Ala Ala 195 200 205

Gln Ala Gln Lys Trp Gly Gly Thr Ala Gly Gln Ala Phe Asp Arg Cys 210 215 220

Tyr His Ser Ser Cys Asp Ser Leu Ser Asn Ile Asn Asp Thr Ala Leu 225 230 235 240

Asp Arg Asn Ser Asp Ala Ala Ala His Ala Ile Trp Thr Leu Ser Ser 245 250 255

## Gly Thr Gly Glu Pro Pro Thr 260

<210> 56

<211> 282

<212> PRT

<213> Homo sapiens

<400> 56

Asp Ala Gln Lys Leu Leu Glu Lys Met Gly Gly Ser Ala Pro Pro Asp 1 5 10 15

Ser Ser Trp Arg Gly Ser Leu Lys Val Pro Tyr Asn Val Gly Pro Gly
20 25 30

Phe Thr Gly Asn Phe Ser Thr Gln Lys Val Lys Met His Ile His Ser 35 40 45

Thr Asn Glu Val Thr Arg Ile Tyr Asn Val Ile Gly Thr Leu Arg Gly 50 55 60

Ala Val Glu Pro Asp Arg Tyr Val Ile Leu Gly Gly His Arg Asp Ser 65 70 75 80

Trp Val Phe Gly Gly Ile Asp Pro Gln Ser Gly Ala Ala Val Val His 85 90 95

Glu Ile Val Arg Ser Phe Gly Thr Leu Lys Lys Glu Gly Trp Arg Pro 100 105 110

Arg Arg Thr Ile Leu Phe Ala Ser Trp Asp Ala Glu Glu Phe Gly Leu 115 120 125

Leu Gly Ser Thr Glu Trp Ala Glu Glu Asn Ser Arg Leu Leu Gln Glu 130 135 140

Arg Gly Val Ala Tyr Ile Asn Ala Asp Ser Ser Ile Glu Gly Asn Tyr 145 150 155 160

Thr Leu Arg Val Asp Cys Thr Pro Leu Met Tyr Ser Leu Val His Asn 165 170 175

Leu Thr Lys Glu Leu Lys Ser Pro Asp Glu Gly Phe Glu Gly Lys Ser 180 185 190

Leu Tyr Glu Ser Trp Thr Lys Lys Ser Pro Ser Pro Glu Phe Ser Gly
195 200 205

Met Pro Arg Ile Ser Lys Leu Gly Ser Gly Asn Asp Phe Glu Val Phe 210 215 220

Phe Gln Arg Leu Gly Ile Ala Ser Gly Arg Ala Arg Tyr Thr Lys Asn 225 230 235 240

Trp Glu Thr Asn Lys Phe Ser Gly Tyr Pro Leu Tyr His Ser Val Tyr

250

Glu Thr Tyr Glu Leu Val Glu Lys Phe Tyr Asp Pro Met Phe Lys Tyr 260 265 270

255

His Leu Thr Val Ala Gln Val Arg Gly Gly 275 280

<210> 57

<211> 282

<212> PRT

<213> Homo sapiens

<400> 57

Asp Ala Glu Ile Leu Leu Arg Tyr Leu Gly Gly Ile Ala Pro Pro Asp 1 5 10 15

Lys Ser Trp Lys Gly Ala Leu Asn Val Ser Tyr Ser Ile Gly Pro Gly
20 25 30

Phe Thr Gly Ser Asp Ser Phe Arg Lys Val Arg Met His Val Tyr Asn 35 40 45

Ile Asn Lys Ile Thr Arg Ile Tyr Asn Val Val Gly Thr Ile Arg Gly
50 55 60

Ser Val Glu Pro Asp Arg Tyr Val Ile Leu Gly Gly His Arg Asp Ser 70 75 80

Trp Val Phe Gly Ala Ile Asp Pro Thr Ser Gly Val Ala Val Leu Gln
85 90 95

Glu Ile Ala Arg Ser Phe Gly Lys Leu Met Ser Lys Gly Trp Arg Pro 100 105 110

Arg Arg Thr Ile Ile Phe Ala Ser Trp Asp Ala Glu Glu Phe Gly Leu 115 120 125

Leu Gly Ser Thr Glu Trp Ala Glu Glu Asn Val Lys Ile Leu Gln Glu 130 135 140

Arg Ser Ile Ala Tyr Ile Asn Ser Asp Ser Ser Ile Glu Gly Asn Tyr 145 150 155 160

Thr Leu Arg Val Asp Cys Thr Pro Leu Leu Tyr Gln Leu Val Tyr Lys 165 170 175

Leu Thr Lys Glu Ile Pro Ser Pro Asp Asp Gly Phe Glu Ser Lys Ser

Leu Tyr Glu Ser Trp Leu Glu Lys Asp Pro Ser Pro Glu Asn Lys Asn 195 200 205

Leu Pro Arg Ile Asn Lys Leu Gly Ser Gly Ser Asp Phe Glu Ala Tyr 210 215 220 Phe Gln Arg Leu Gly Ile Ala Ser Gly Arg Ala Arg Tyr Thr Lys Asn 225 230 235 240

Lys Lys Thr Asp Lys Tyr Ser Ser Tyr Pro Val Tyr His Thr Ile Tyr 245 250 255

Glu Thr Phe Glu Leu Val Glu Lys Phe Tyr Asp Pro Thr Phe Lys Lys 260 265 270

Gln Leu Ser Val Ala Gln Leu Arg Gly Ala 275 280

<210> 58

<211> 283

<212> PRT

<213> Homo sapiens

<400> 58

Arg Asp Leu Cys Asn Leu Asn Gly Thr Leu Ala Pro Ala Thr Trp  $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$ 

Gln Gly Ala Leu Gly Cys His Tyr Arg Leu Gly Pro Gly Phe Arg Pro 20 25 30

Asp Gly Asp Phe Pro Ala Asp Ser Gln Val Asn Val Ser Val Tyr Asn 35 40 45

Arg Leu Glu Leu Arg Asn Ser Ser Asn Val Leu Gly Ile Ile Arg Gly 50 55 60

Ala Val Glu Pro Asp Arg Tyr Val Leu Tyr Gly Asn His Arg Asp Ser 65 70 75 80

Trp Val His Gly Ala Val Asp Pro Ser Ser Gly Thr Ala Val Leu Leu
85 90 95

Glu Leu Ser Arg Val Leu Gly Thr Leu Leu Lys Lys Gly Thr Trp Arg 100 105 110

Pro Arg Arg Ser Ile Val Phe Ala Ser Trp Gly Ala Glu Glu Phe Gly 115 120 125

Leu Ile Gly Ser Thr Glu Phe Thr Glu Glu Phe Phe Asn Lys Leu Gln 130 135 140

Glu Arg Thr Val Ala Tyr Ile Asn Val Asp Ile Ser Val Phe Ala Asn 145 150 155 160

Ala Thr Leu Arg Val Gln Gly Thr Pro Pro Val Gln Ser Val Val Phe 165 170 175

Ser Ala Thr Lys Glu Ile Arg Ser Pro Gly Pro Gly Asp Leu Ser Ile 180 185 190 Tyr Asp Asn Trp Ile Arg Tyr Phe Asn Arg Ser Ser Pro Val Tyr Gly
195 200 205

Leu Val Pro Ser Leu Gly Ser Leu Gly Ala Gly Ser Asp Tyr Ala Pro 210 215 220

Phe Val His Phe Leu Gly Ile Ser Ser Met Asp Ile Ala Tyr Thr Tyr 225 230 235 240

Asp Arg Ser Lys Thr Ser Ala Arg Ile Tyr Pro Thr Tyr His Thr Ala 245 250 255

Phe Asp Thr Phe Asp Tyr Val Asp Lys Phe Leu Asp Pro Gly Phe Ser 260 265 270

Ser His Gln Ala Val Ala Arg Thr Ala Gly Ser 275 280

<210> 59

<211> 259

<212> PRT

<213> Homo sapiens

<400> 59

Ser Pro His Thr Gly Ile Gln Glu Tyr Gln Asp Gly Val Pro Lys Ile 5 10 15

Pro Thr Ala Cys Ile Thr Val Glu Asp Ala Glu Met Met Ser Arg Met 20 25 30

Ala Ser His Gly Ile Lys Ile Val Ile Gln Leu Lys Met Gly Ala Lys 35 40 45

Thr Tyr Pro Asp Thr Asp Ser Phe Asn Thr Val Ala Glu Ile Thr Gly 50 55 60

Ser Lys Tyr Pro Glu Gln Val Val Leu Val Ser Gly His Leu Asp Ser 65 70 75 80

Trp Asp Val Gly Gln Gly Ala Met Asp Asp Gly Gly Gly Ala Phe Ile 85 90 95

Ser Trp Glu Ala Leu Ser Leu Ile Lys Asp Leu Gly Leu Arg Pro Lys 100 105 110

Arg Thr Leu Arg Leu Val Leu Trp Thr Ala Glu Glu Gln Gly Gly Val 115 120 125

Gly Ala Phe Gln Tyr Tyr Gln Leu His Lys Val Asn Ile Ser Asn Tyr 130 135 140

Ser Leu Val Met Glu Ser Asp Ala Gly Thr Phe Leu Pro Thr Gly Leu 145 150 155 160

Gln Phe Thr Gly Ser Glu Lys Ala Arg Ala Ile Met Glu Glu Val Met

				165					1/0					1/5	
Ser	Leu	Leu	Gln 180	Pro	Leu	Asn	Ile	Thr 185	Gln	Val	Leu	Ser	His 190	Gly	Glu
Gly	Thr	Asp 195	Ile	Asn	Phe	Trp	Ile 200	Gln	Ala	Gly	Val	Pro 205	Gly	Ala	Ser

Leu Leu Asp Asp Leu Tyr Lys Tyr Phe Phe Phe His His Ser His Gly 210 215 220

Asp Thr Met Thr Val Met Asp Pro Lys Gln Met Asn Val Ala Ala 225 230 235 240

Val Trp Ala Val Val Ser Tyr Val Val Ala Asp Met Glu Glu Met Leu 245 250 255

Pro Arg Ser